



Blast 2 Sequences results

[PubMed](#)[Entrez](#)[BLAST](#)[OMIM](#)[Taxonomy](#)[Structure](#)

BLAST 2 SEQUENCES RESULTS VERSION BLASTN 2.2.18 [Mar-02-2008]

Match: Mismatch: gap open: gap extension:
x_dropoff: expect: wordsize: Filter View option
Masking character option Masking color option
 Show CDS translation

Sequence 1: gi|184371|Human hepsin mRNA, complete cds

Length = 1783

Sequence 2: gi|14602458|Homo sapiens transmembrane protease, serine 2 (TMPRSS2), mRNA
Length = 3226

No significant similarity was found

CPU time: 0.05 user secs. 0.03 sys. secs 0.08 total secs.



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BLAST 2 SEQUENCES RESULTS VERSION BLASTN 2.2.18 [Mar-02-2008]

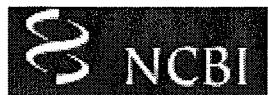
Match: 1 Mismatch: -2 gap open: 5 gap extension: 2
x_dropoff: 0 expect: 10.000 wordsize: 11 Filter View option Standard
Masking character option X for protein, n for nucleotide Masking color option Black
 Show CDS translation

Sequence 1: gi|184371|Human hepsin mRNA, complete cds
Length = 1783

Sequence 2: gi|13173470|Homo sapiens transmembrane protease, serine 3 (TMPRSS3), transcript variant A, mRNA >gi|12246823|dbj|AB038157.1| Homo sapiens TMPRSS3a mRNA for serine protease, complete cds
Length = 2468

No significant similarity was found

CPU time: 0.05 user secs. 0.04 sys. secs 0.09 total secs.



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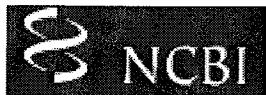
Match: 1 Mismatch: -2 gap open: 5 gap extension: 2
x_dropoff: 0 expect: 10.0000 wordsize: 11 Filter View option Standard
Masking character option X for protein, n for nucleotide Masking color option Black
 Show CDS translation

Sequence 1: gi|184371|Human hepsin mRNA, complete cds
Length = 1783

Sequence 2: gi|145701031|Homo sapiens transmembrane protease, serine 4 (TMPRSS4), transcript variant 1, mRNA
Length = 3549

No significant similarity was found

CPU time: 0.05 user secs. 0.05 sys. secs 0.10 total secs.



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BLAST 2 SEQUENCES RESULTS VERSION BLASTN 2.2.18 [Mar-02-2008]

Match: 1 Mismatch: -2 gap open: 5 gap extension: 2

x_dropoff: 0 expect: 10.0000 wordsize: 11 Filter View option Standard

Masking character option X for protein, n for nucleotide Masking color option Black

Show CDS translation

Sequence 1: gi|184371|Human hepsin mRNA, complete cds

Length = 1783

Sequence 2: gi|170932491|Homo sapiens transmembrane protease, serine 5 (spinesin) (TMPRSS5), mRNA

Length = 2233

No significant similarity was found

CPU time: 0.04 user secs. 0.04 sys. secs 0.08 total secs.



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BLAST 2 SEQUENCES RESULTS VERSION BLASTN 2.2.18 [Mar-02-2008]

Match: 1 Mismatch: -2 gap open: 5 gap extension: 2

x_dropoff: 0 expect: 10.0000 wordsize: 11 Filter View option Standard

Masking character option X for protein, n for nucleotide Masking color option Black

Show CDS translation

Sequence 1: gi|184371|Human hepsin mRNA, complete cds

Length = 1783

Sequence 2: gi|9719385|Homo sapiens enteropeptidase (PRSS7) gene, partial cds

Length = 224

No significant similarity was found

CPU time: 0.05 user secs. 0.06 sys. secs 0.11 total secs.



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BLAST 2 SEQUENCES RESULTS VERSION BLASTN 2.2.18 [Mar-02-2008]

Match: 1 Mismatch: -2 gap open: 5 gap extension: 2
x_dropoff: 0 expect: 10.000 wordsize: 11 Filter View option Standard
Masking character option X for protein, n for nucleotide Masking color option Black
 Show CDS translation

Sequence 1: gi|184371|Human hepsin mRNA, complete cds

Length = 1783

Sequence 2: gi|116256362|Homo sapiens transmembrane protease, serine 13 (TMPRSS13), mRNA
Length = 3391

No significant similarity was found

CPU time: 0.05 user secs. 0.04 sys. secs 0.09 total secs.